

Amendments to the Claims:

This listing of claims will replace all prior versions, and listings, of claims in the application:

Listing of Claims:

Claims 1-26 (canceled)

27. (withdrawn) A method of preparing a sialyl Lewis x determinant, the method comprising contacting a murine Fuc-TVII enzyme with a GDP-fucose donor substrate and a sialyl-N-acetyl-lactosamine acceptor substrate in an enzyme bioreactor under conditions that allow the addition of an α 1,3 linked fucose to the sialyl-N-acetyl-lactosamine acceptor substrate, wherein the murine Fuc-TVII enzyme comprises a catalytic domain that is encoded by a nucleic acid sequence segment that is identical to a polynucleotide that is amplified using murine mRNA or cDNA as a template by a 5' primer as shown in SEQ ID NO:3 and a 3' primer as shown in SEQ ID NO:4.

28. (withdrawn) The method of claim 27, wherein the Fuc-TVII enzyme is in solution.

29. (withdrawn) The method of claim 27, wherein the Fuc-TVII enzyme is immobilized on a solid phase matrix.

30. (withdrawn) The method of claim 27, wherein the Fuc-TVII enzyme is a recombinant enzyme.

31. (withdrawn) The method of claim 20, wherein the Fuc-TVII enzyme is produced in a mammalian host cell.

32. (withdrawn) The method of claim 20, wherein the Fuc-TVII enzyme is produced in a baculovirus host.

33. (withdrawn) The method of claim 27, wherein the sialyl-N-acetyl-lactosamine acceptor is on a glycoprotein.

34. (withdrawn) The method of claim 27, wherein the sialyl-N-acetyl-lactosamine acceptor is on a glycolipid.

35. (withdrawn) The method of claim 27, wherein the sialyl-N-acetyl-lactosamine acceptor is a free oligosaccharide.

36. (canceled)

37. (previously presented) A murine Fuc-TVII enzyme comprising a catalytic domain that is encoded by a nucleic acid sequence segment that is identical to a polynucleotide that is amplified using murine mRNA or cDNA as a template by a 5' primer as shown in SEQ ID NO:3 and a 3' primer as shown in SEQ ID NO:4.

38. (previously presented) The murine Fuc-TVII enzyme of claim 37, wherein the catalytic domain is encoded by a nucleic acid segment consisting of residue 2194 to residue 3085 of SEQ ID NO: 1.